

WHAT IS CLAIMED IS:

- 1 1. A method for obtaining a cell-specific binding molecule useful for
2 increasing uptake or specificity of a genetic vaccine to a target cell, the method comprising:
3 creating a library of recombinant polynucleotides that by recombining a
4 nucleic acid that encodes a polypeptide that comprises a nucleic acid binding domain and a
5 nucleic acid that encodes a polypeptide that comprises a cell-specific binding domain; and
6 screening the library to identify a recombinant polynucleotide that
7 encodes a binding molecule that can bind to a nucleic acid and to a cell-specific receptor.
- 1 2. A method for obtaining a cell-specific binding moiety useful for
2 increasing uptake or specificity of a genetic vaccine to a target cell, the method comprising:
3 (1) recombining at least first and second forms of a nucleic acid which
4 comprises a polynucleotide that encodes a nucleic acid binding domain and at least first and
5 second forms of a nucleic acid which comprises a cell-specific ligand that specifically binds
6 to a protein on the surface of a cell of interest, wherein the first and second forms differ from
7 each other in two or more nucleotides, to produce a library of recombinant binding moiety-
8 encoding nucleic acids;
9 (2) transfecting into a population of host cells a library of vectors, each
10 of which comprises: a) a binding site specific for the nucleic acid binding domain and 2) a
11 member of the library of recombinant binding moiety-encoding nucleic acids, wherein the
12 recombinant binding moiety is expressed and binds to the binding site to form a vector-
13 binding moiety complex;
14 (3) lysing the host cells under conditions that do not disrupt binding of
15 the vector-binding moiety complex;
16 (4) contacting the vector-binding moiety complex with a target cell of
17 interest; and
18 (5) identifying target cells that contain a vector and isolating the
19 optimized recombinant cell-specific binding moiety nucleic acids from these target cells.
- 1 3. The method of claim 2, wherein the method further comprises:

2 (6) recombining at least one optimized recombinant binding moiety,
3 encoding nucleic acid with a further form of the polynucleotide that encodes a nucleic acid
4 binding domain and/or a further form of the polynucleotide that encodes a cell-specific
5 ligand, which are the same or different from the first and second forms, to produce a further
6 library of recombinant binding moiety-encoding nucleic acids;

(7) transfected into a population of host cells a library of vectors that comprise: a) a binding site specific for the nucleic acid binding domain and 2) the recombinant binding moiety-encoding nucleic acids, wherein the recombinant binding moiety is expressed and binds to the binding site to form a vector-binding moiety complex

(8) lysing the host cells under conditions that do not disrupt binding of the vector-binding moiety complex;

(9) contacting the vector-binding moiety complex with a target cell of interest and identifying target cells that contain the vector; and

(10) isolating the optimized recombinant binding moiety nucleic acids from the target cells which contain the vector; and

(11) repeating (6) through (10), as necessary, to obtain a further optimized cell-specific binding moiety useful for increasing uptake or specificity of a genetic vaccine vector to a target cell.

1 4. The method of claim 2, wherein the method further comprises
2 identifying cell-specific binding moieties that result in the highest efficiency in transfecting
3 the target cells.

1 5. The method of claim 2, wherein the nucleic acid binding domain is a
2 DNA binding domain derived from a protein selected from the group consisting of a
3 transcriptional regulator, a polypeptide involved in DNA replication or recombination, a
4 repressor, a histone, a protamine, an *E. coli* CAP protein, myc, a protein having a leucine
5 zipper, a protein having a DNA binding basic domain, a protein having a POU domain, a
6 protein having a zinc finger, and a protein having a Cys₃His box.

1 6. The method of claim 2, wherein the nucleic acid binding domain is an
2 RNA binding domain derived from a protein selected from the group consisting of HIV *tat*
3 and HIV *rev*.

1 7. The method of claim 2, wherein the target cell of interest is selected
2 from the group consisting of muscle cells, monocytes, dendritic cells, B cells, Langerhans
3 cells, keratinocytes, and M-cells.

1 8. The method of claim 7, wherein the cell of interest is a professional
2 antigen presenting cell.

1 9. The method of claim 8, wherein the antigen presenting cell is a dendritic
2 cell, a monocyte/macrophage, a B cell, or a Langerhans cell.

1 10. The method of claim 8, wherein the cell-specific ligand comprises a
2 polypeptide selected from the group consisting of CD2, CD28, CTLA-4, CD40 ligand,
3 fibrinogen, ICAM-1, Fc portion of immunoglobulin G, and a bacterial enterotoxin, or a
4 subunit thereof.

1 11. The method of claim 2, wherein the target cell of interest is a human
2 cell.

1 12. The method of claim 2, wherein the target cells that contain the vector
2 are identified by selecting for expression of a selectable marker contained in the vector.

1 13. The method of claim 2, wherein the optimized recombinant binding
2 moiety-encoding nucleic acid comprises a genetic vaccine vector.

1 14. A cell-specific recombinant binding moiety produced by expressing in a
2 host cell an optimized recombinant binding moiety-encoding nucleic acid obtained by the
3 method of claim 2.

1 15. A genetic vaccine that comprises a cell-specific recombinant binding
2 moiety of claim 14;

1 16. A genetic vaccine that comprises an optimized recombinant binding
2 moiety-encoding nucleic acid obtained by the method of claim 2;

1 17. A genetic vaccine that comprises:
2 a) an optimized recombinant binding moiety that comprises a nucleic
3 acid binding domain and a cell-specific ligand, and
4 b) a polynucleotide sequence that comprises a binding site, wherein the
5 nucleic acid binding domain is capable of specifically binding to the binding site.

1 18. A method for obtaining an optimized cell-specific binding moiety useful
2 for increasing uptake, efficacy, or specificity of a genetic vaccine for a target cell, the
3 method comprising:

4 (1) recombining at least first and second forms of a nucleic acid that
5 comprises a polynucleotide which encodes a non-toxic receptor binding moiety of an
6 enterotoxin, wherein the first and second forms differ from each other in two or more
7 nucleotides, to produce a library of recombinant nucleic acids;
8 (2) transfecting vectors that contain the library of nucleic acids into a
9 population of host cells, wherein the nucleic acids are expressed to form recombinant cell-
10 specific binding moiety polypeptides;
11 (3) contacting the recombinant cell-specific binding moiety
12 polypeptides with a cell surface receptor of a target cell; and
13 (4) determining which recombinant cell-specific binding moiety
14 polypeptides exhibit enhanced ability to bind to the target cell.

1 19. The method of claim 18, wherein the cell surface receptor is present on
2 the surface of a target cell.

1 20. The method of claim 18, wherein the cell surface receptor is G_{M1}.

1 21. The method of claim 18, wherein the host cell is a *V. cholerae* cell
2 which is incapable of expressing CT-A.

1 22. A method for enhancing uptake of a genetic vaccine vector by a target
2 cell, the method comprising coating the genetic vaccine vector with an optimized
3 recombinant cell-specific binding moiety produced by the method of claim 18,

1 23. The method of claim 18, wherein the recombinant cell-specific binding
2 moieties are expressed as a fusion protein on the surface of a replicable genetic package.

1 24. A method of obtaining a genetic vaccine component that confers upon a
2 vector an enhanced ability to enter an antigen-presenting cell, the method comprising:
3 creating a library of recombinant nucleic acids by subjecting to
4 recombination at least two forms of a polynucleotide;
5 contacting a library of vectors, each of which comprises a member of
6 the library of recombinant nucleic acids, with a population of antigen-presenting or antigen-
7 processing cells; and
8 determining the percentage of cells in the population that contain the
9 vector.

1 25. The method of claim 24, wherein the antigen-presenting or antigen-
2 processing cells are selected from the group consisting of B cells, monocytes/macrophages,
3 dendritic cells, Langerhans cells, keratinocytes, and muscle cells.

1 26. The method of claim 25, wherein the cells are B cells which are
2 obtained from a B cell line.

1 27. The method of claim 24, wherein the screening is conducted *in vivo* and
2 the cells are monkey cells or mouse cells.

1 28. The method of claim 24, wherein the method further comprises:

2 culturing the cells for a predetermined time after contacting the cells
3 with the library of vectors;

washing the cells after the contacting step to remove vectors that did not enter an antigen-presenting cell; and

isolating the vectors from the cells that contain a vector.

1 29. The method of claim 24, wherein the cells that contain a vector are
2 identified by:

5 co-culturing the cultures of antigen-presenting cells with T lymphocytes
6 obtained from the same individual as the antigen-presenting cells; and

7 identifying cultures in which a T lymphocyte response is induced.

1 30. The method of claim 29, wherein the T lymphocyte response is selected
2 from the group consisting of increased T lymphocyte proliferation, increased T lymphocyte-
3 mediated cytolytic activity against a target cell, and increased cytokine production.

1 31. The method of claim 24, wherein the vector is a replicable genetic
2 package and the recombinant nucleic acids are expressed as a fusion protein which is
3 displayed on the surface of the replicable genetic package.

32. The method of claim 31, wherein the replicable genetic package is a
bacteriophage.

1 33. A method of obtaining a genetic vaccine component that confers upon a
2 vector an enhanced ability to enter cell or tissue when administered to a mammal by a
3 desired administration protocol, the method comprising:

4 creating a library of recombinant nucleic acids by subjecting to
5 recombination at least two forms of a polynucleotide;

6 / administering to a mammal a library of vectors, each of which
7 comprises a member of the library of recombinant nucleic acids, into a mammal;

8 obtaining target cells or tissues from the mammal;
9 identifying target cells or tissues that contain a vector, and
10 recovering vectors from the identified target cells or tissues.

1 34. The method of claim 33, wherein the target cells are lymphatic cells.

1 35. The method of claim 33, wherein the administering is by oral ingestion,
2 inhalation, injection, or topical application to skin or mucous membrane.

1 36. The method of claim 33, wherein the vector is a replicable genetic
2 package and the recombinant nucleic acids are expressed as a fusion protein which is
3 displayed on the surface of the replicable genetic package.

1 37. A method for evolving a vaccine delivery vehicle to obtain an optimized
2 delivery vehicle having enhanced ability to enter a selected mammalian tissue upon
3 administration to a mammal, the method comprising:

4 (1) recombining members of a pool of polynucleotides to produce a
5 library of recombinant polynucleotides;

6 (2) administering to a test animal a library of replicable genetic
7 packages, each of which comprises a member of the library of recombinant polynucleotides
8 operably linked to a polynucleotide that encodes a display polypeptide, wherein the
9 recombinant polynucleotide and the display polypeptide are expressed as a fusion protein
10 which is displayed on the surface of the replicable genetic package; and

11 (3) recovering replicable genetic packages that are present in the
12 selected tissue of the test animal at a suitable time after administration, wherein recovered
13 replicable genetic packages have enhanced ability to enter the selected mammalian tissue
14 upon administration to the mammal.

1 38. The method of claim 37, wherein the method further comprises:

2 (4) recombining a nucleic acid that comprises at least one recombinant
3 polynucleotide obtained from a replicable genetic package recovered from the selected tissue

4 with a further pool of polynucleotides to produce a further library of recombinant
5 polynucleotides;

6 (5) administering to a test animal a library of replicable genetic
7 packages, each of which comprises a member of the further library of recombinant
8 polynucleotides operably linked to a polynucleotide that encodes a display polypeptide,
9 wherein the recombinant polynucleotide and the display polypeptide are expressed as a
10 fusion protein which is displayed on the surface of the replicable genetic package;

11 (6) recovering replicable genetic packages that are present in the
12 selected tissue of the test animal at a suitable time after administration; and

13 (7) repeating (4) through (6), as necessary, to obtain a further optimized
14 recombinant delivery vehicle that exhibits further enhanced ability to enter a selected
15 mammalian tissue upon administration to a mammal.

1 39. The method of claim 37, wherein the replicable genetic package is a
2 bacteriophage.

1 40. The method of claim 39, wherein the bacteriophage is M13.

1 41. The method of claim 40, wherein the polynucleotide which encodes a
2 display polypeptide is selected from the group consisting of gene III and gene VIII.

1 42. The method of claim 37, wherein the selected mammalian tissue is the
2 bloodstream and the administration is by inhalation.

1 43. The method of claim 37, wherein the administration is intravenous and
2 the selected mammalian tissue is selected from the group consisting of lymph node and
3 spleen.

1 44. A method for evolving a vaccine delivery vehicle to obtain an optimized
2 delivery vehicle having enhanced specificity for antigen-presenting cells, the method
3 comprising:

(1) recombining members of a pool of polynucleotides to produce a library of recombinant polynucleotides;

6 (2) producing a library of replicable genetic packages, each of which
7 comprises a member of the library of recombinant polynucleotides operably linked to a
8 polynucleotide that encodes a display polypeptide, wherein the recombinant polynucleotide
9 and the display polypeptide are expressed as a fusion protein which is displayed on
10 the surface of the replicable genetic package;

11 (3) contacting the library of recombinant replicable genetic packages
12 with a non-APC to remove replicable genetic packages that display non-APC-specific fusion
13 polypeptides; and

14 (4) contacting the recombinant replicable genetic packages that did not
15 bind to the non-APC with an APC and recovering those that bind to the APC, wherein the
16 recovered replicable genetic packages are capable of specifically binding to APCs.

1 45. The method of claim 44, wherein the method further comprises the steps
2 of:

3 (5) recombining a nucleic acid which comprises at least one
4 recombinant polynucleotide obtained from a replicable genetic package that is capable of
5 specifically binding to APCs with a further pool of polynucleotides to produce a further
6 library of recombinant polynucleotides;

7 (6) producing a further library of recombinant replicable genetic
8 packages, each of which comprises a member of the library of recombinant polynucleotides
9 operably linked to a polynucleotide that encodes a display polypeptide, wherein the
10 recombinant polynucleotide and the display polypeptide are expressed as a fusion protein
11 which is displayed on the surface of the replicable genetic package;

(7) contacting the further library of recombinant replicable genetic packages with a non-APC to remove those that display non-APC-specific fusion polypeptides; and

15 (8) contacting the recombinant replicable genetic packages which did
16 not bind to the non-APC with an APC and recovering replicable genetic packages which
17 bind to the APC, wherein the recovered replicable genetic packages are capable of
18 specifically binding to APCs; and

19 (9) repeating (5) through (8), as necessary, to obtain a further optimized
20 recombinant delivery vehicle which exhibits further enhanced specificity for antigen-
21 presenting cells.

46. A method for evolving a vaccine delivery vehicle to obtain an optimized
delivery vehicle having enhanced ability to enter a target cell, the method comprising:

(1) recombining at least first and second forms of a nucleic acid which encodes an invasin polypeptide, wherein the first and second forms differ from each other in two or more nucleotides, to produce a library of recombinant invasin nucleic acids;

10 (3) contacting the library of recombinant bacteriophage with a
11 population of target cells;

12 (4) removing unbound phage and phage which is bound to the surface
13 of the target cells; and

(5) recovering phage which are present within the target cells, wherein the recovered phage are enriched for phage that have enhanced ability to enter the target cells.

1 47. The method of claim 46, wherein the method further comprises:

2 (6) recombining a nucleic acid which comprises at least one
3 recombinant invasin nucleic acid obtained from a bacteriophage which is recovered from a
4 target cell with a further pool of polynucleotides to produce a further library of recombinant
5 invasin polynucleotides;

(7) producing a further library of recombinant bacteriophage, each of which displays on the bacteriophage surface a fusion polypeptide encoded by a chimeric gene that comprises a recombinant invasin nucleic acid operably linked to a polynucleotide that encodes a display polypeptide;

(8) contacting the library of recombinant bacteriophage with a population of target cells;

1 50. The method of claim 46, wherein the target cell is an APC.

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